152

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.

190 66 175 118-180-

 $\frac{256}{257}$   $\frac{257}{275}$   $\frac{257}{275}$ 

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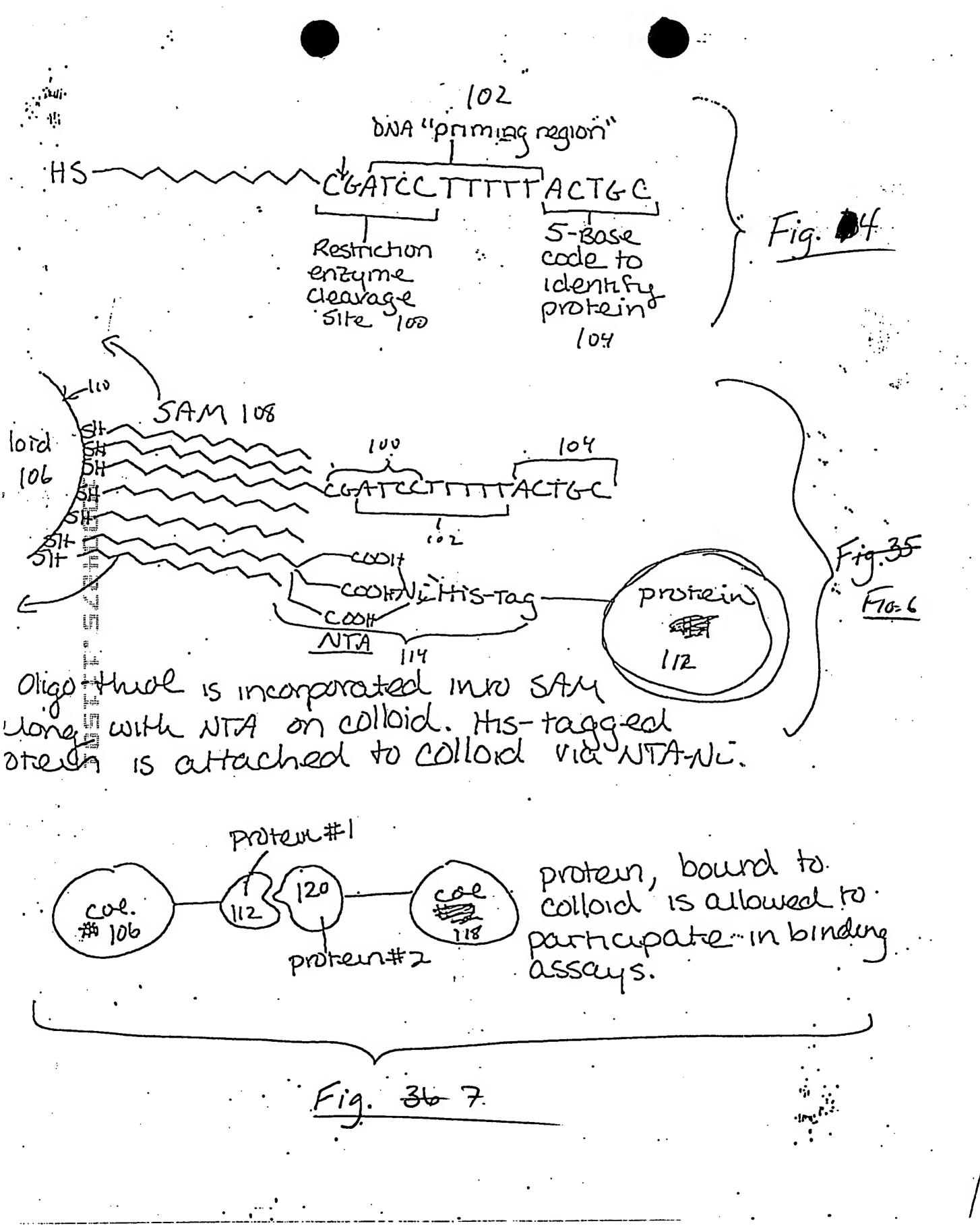
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Complementary DNA to "DNA priming region" sequence using standard PCR methods: STAGGAAAA Sequence complementary oligo: 122 match up resulting sequence bata with records. Kept that connect protein identify to sequence: ACTGC = protein # 120



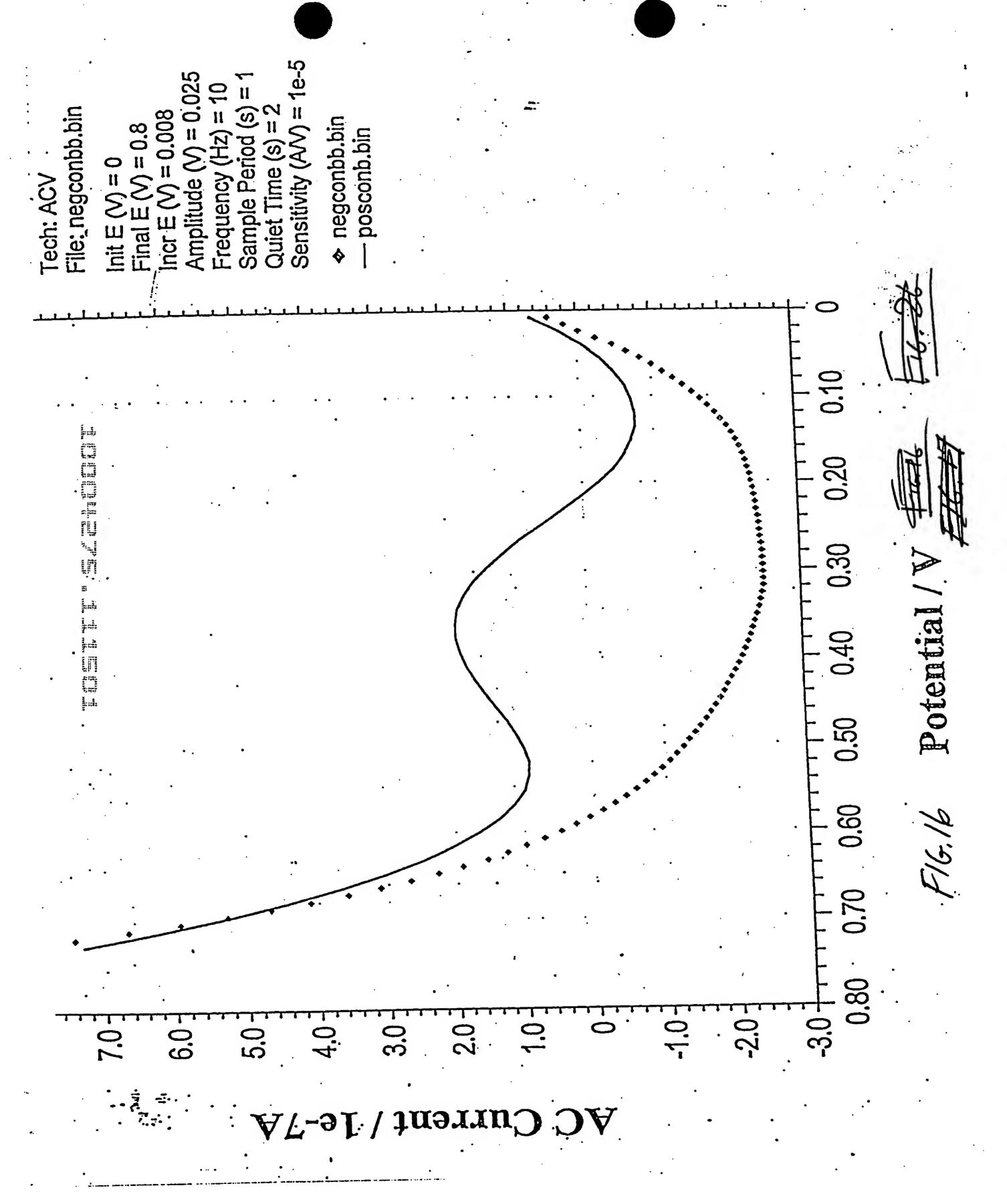
To uncover the identity of proteen after the assays are completted, cleave the DNA portion of the DNA-thurk by addition of a restriction enzyme: Restriction enzyme cleaves DNA.

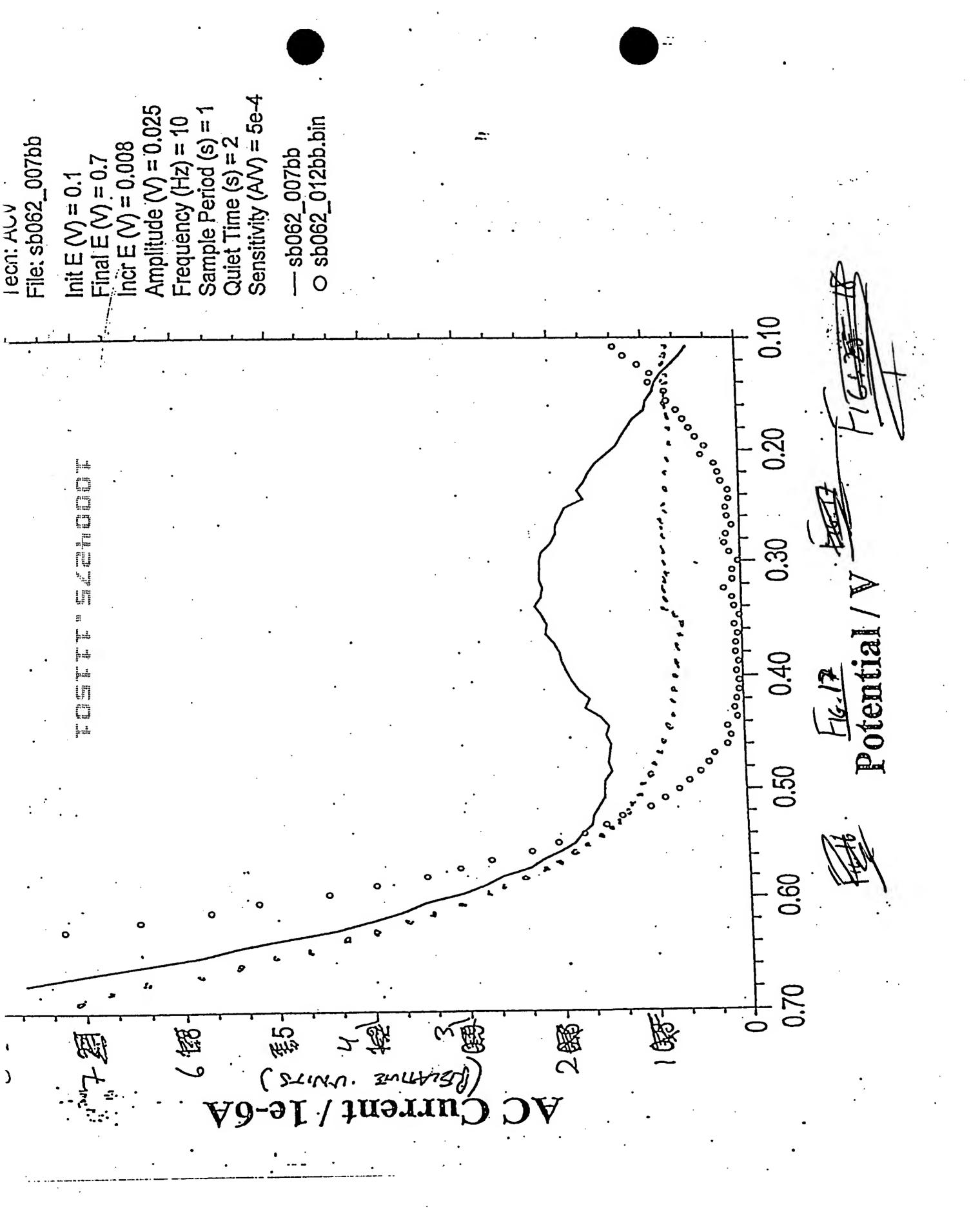
Lat specific recognition site colloid **举118** NTA-Itis linkage protein) H16,8 102 104

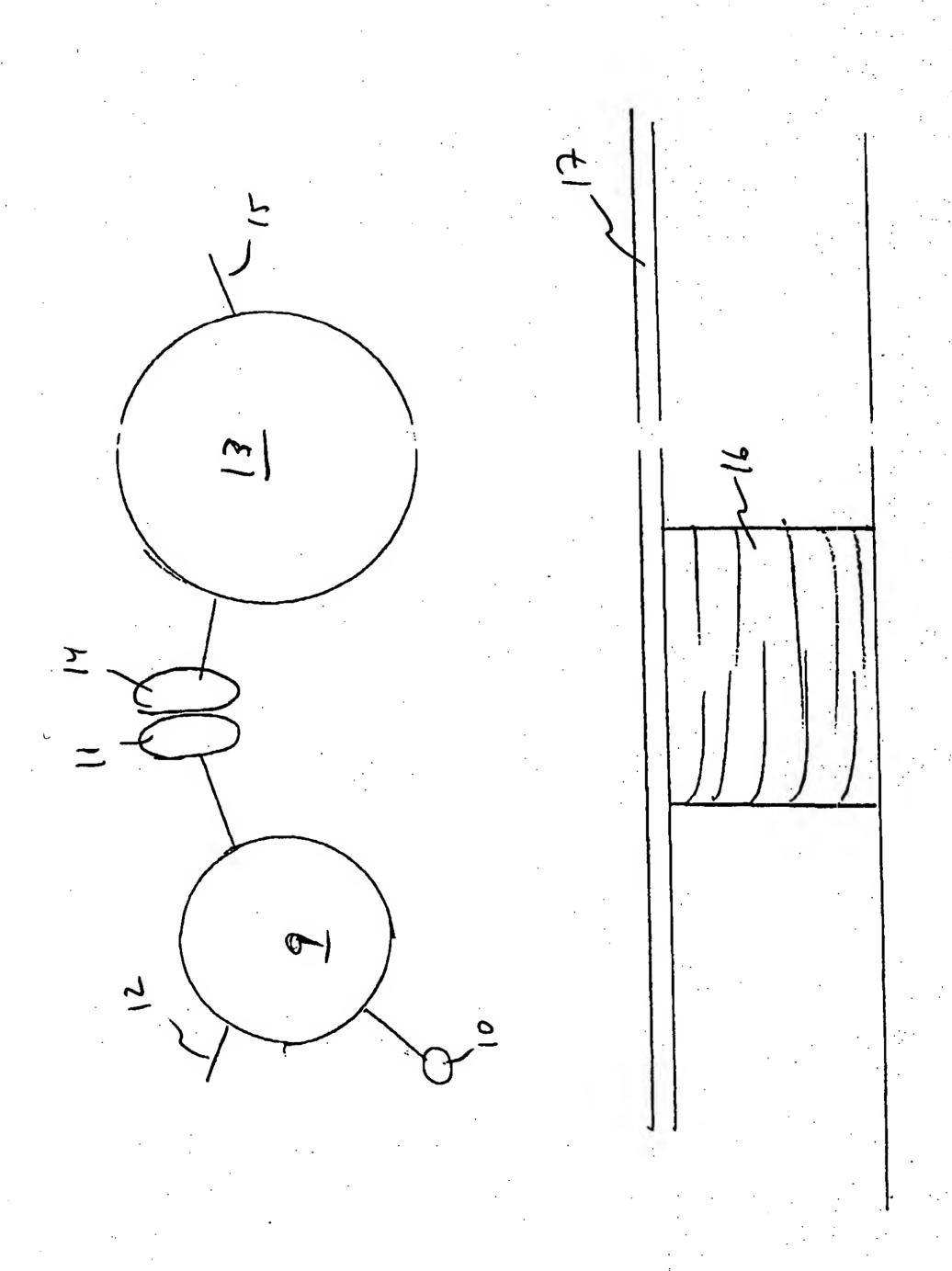
chematic #2: wique identifying sequence colloid 28 DWA throe is incorporated into samon colloid along with NTA+ protein is attached via Itis-tag coffords bearing proteins or small molecules are allowed to interact. Birding of protein XI to small molecule Y attouts brings their DNA tags into close proximity/colloid 134

2. Complementary sequences to fanction DNA Hags are added + allowed to bind. collow colloid. Y 130 Remove b Single-strainded Dutaise is added to remove (or ""chew up" any 5H non-hybridized DNA. Result 136 GACTG-TCATCG CTG-ACAG-TAGE Collex Colloid 128 x 124 132 126 : Flb. 13 134 Colleid

Complementary DNA is denarured and -TGACTG-TCATCG CIGALAGTI 132 130 Resulting sequence contains the unique DNA codes of the two binding partners, x +c 36 AGTAGC unique sequence unque protein sequence ( 5,5EC,ES 137) Protein X (50ECLES 126) Protein X + Protein / must be binding. partners.







F16.18

